

PCT

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RAW SEQUENCE LISTING
                                                              DATE: 11/06/2003
                     PATENT APPLICATION: US/10/089,473A
                                                              TIME: 12:15:54
                     Input Set : A:\PTO.KD.txt
                     Output Set: N:\CRF4\11062003\J089473A.raw
      3 <110> APPLICANT: CHUNG, Kwang-Hoe
              KIM, Doo-Sik
              KOH, You-Seok
              SOHN, Young-Dong
              YOU, Weon-Kyoo
              JANG, Yang-Soo
              HUH, Chin-Kyu
     11 <120> TITLE OF INVENTION: NOVEL PROTEIN DERIVED FROM AGKISTRODON SAXATILIS EMELIANOV
AND PROCESS
              FOR PREPARING THE SAME
     14 <130> FILE REFERENCE: 0136/0K432US0
     16 <140> CURRENT APPLICATION NUMBER: US 10/089,473A
C--> 17 <141> CURRENT FILING DATE: 2003-10-15
                                                            ENTERED
     19 <160> NUMBER OF SEQ ID NOS: 10
     21 <170> SOFTWARE: PatentIn version 3.1
     23 <210> SEQ ID NO: 1
     24 <211> LENGTH: 73
     25 <212> TYPE: PRT
     26 <213> ORGANISM: Agkistrodon saxatilis emelianov
     28 <400> SEQUENCE: 1
    -30 Glu Ala Gly Glu Glu Cys Asp Cys Gly Ala Pro Ala Asn Pro Cys Cys
     31 1
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     34 Asp Ala Ala Thr Cys Lys Leu Arg Pro Gly Ala Gln Cys Ala Glu Gly
                                        25
     38 Leu Cys Cys Asp Gln Cys Arg Phe Met Lys Glu Gly Thr Ile Cys Arg
                35
                                    40
     42 Met Ala Arg Gly Asp Asp Met Asp Asp Tyr Cys Asn Gly Ile Ser Ala
     46 Gly Cys Pro Arg Asn Pro Phe His Ala
     47 65
     50 <210> SEQ ID NO: 2
     51 <211> LENGTH: 213
     52 <212> TYPE: DNA
     53 <213> ORGANISM: Agkistrodon saxatilis emelianov
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58 ctgagaccag gggcgcagtg tgcagaagga ctgtgttgtg accagtgcag atttatgaaa

60 gaaggaacaa tatgccggat ggcaaggggt gatgacatgg atgattactg caatggcata

55 <400> SEQUENCE: 2

65 <210> SEQ ID NO: 3 66 <211> LENGTH: 20 67 <212> TYPE: DNA

70 <220> FEATURE:

62 tetgetgget gteccagaaa teeetteeat gee

68 <213> ORGANISM: Artificial Sequence

5

6

9

12

35

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180

213

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71 <223> OTHER INFORMATION: primer
     73 <220> FEATURE:
     74 <221> NAME/KEY: misc feature
     75 <222> LOCATION: (1)..(20)
     76 <223> OTHER INFORMATION: where n may be a or g or c or t/u, other, or unknown
     79 <400> SEQUENCE: 3
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     83 <210> SEQ ID NO: 4
     84 <211> LENGTH: 20
     85 <212> TYPE: DNA
     86 <213> ORGANISM: Artificial Sequence
     88 <220> FEATURE:
     89 <223> OTHER INFORMATION: primer
     91 <400> SEQUENCE: 4
     92 ggcatggaag ggatttctgg
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     95 <210> SEQ ID NO: 5
     96 <211> LENGTH: 33
     97 <212> TYPE: DNA
     98 <213> ORGANISM: Artificial Sequence
     100 <220> FEATURE:
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     103 <400> SEQUENCE: 5
     104 ccgctcgaga aaagagaggc cggagaagaa tgt
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     109 <212> TYPE: DNA
     110 <213> ORGANISM: Artificial Sequence
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    115 <400> SEQUENCE: 6
    116 cggaattctc attaggcatg gaaggga
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     122 <213> ORGANISM: Artificial Sequence
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     133 <210> SEO ID NO: 8
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    136 <213> ORGANISM: Artificial Sequence
    138 <220> FEATURE:
    139 <223> OTHER INFORMATION: oligopeptide
    141 <400> SEQUENCE: 8
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147 <210> SEQ ID NO: 9 148 <211> LENGTH: 12 149 <212> TYPE: PRT 150 <213> ORGANISM: Agkistrodon saxatilis emelianov 152 <400> SEQUENCE: 9 154 Gly Glu Glu Cys Asp Cys Gly Ala Pro Ala Asn Pro 158 <210> SEQ ID NO: 10 159 <211> LENGTH: 73 160 <212> TYPE: PRT 161 <213> ORGANISM: Agkistrodon halys brevicaudus 163 <400> SEQUENCE: 10 165 Glu Ala Gly Glu Cys Asp Cys Gly Ser Pro Gly Asn Pro Cys Cys 169 Asp Ala Ala Thr Cys Lys Leu Arg Gln Gly Ala Gln Cys Ala Glu Gly 20 173 Leu Cys Cys Asp Gln Cys Arg Phe Met Lys Glu Gly Thr Ile Cys Arg 177 Arg Ala Arg Gly Asp Asp Leu Asp Asp Tyr Cys Asn Gly Ile Ser Ala 55 181 Gly Cys Pro Arg Asn Pro Phe His Ala 182 65

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/089,473A

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:3; N Pos. 3